

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/522,592
Source: PT/10
Date Processed by STIC: 2/22/05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
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Revised 01/24/05



PCT

RAW SEQUENCE LISTING

DATE: 02/22/2005

PATENT APPLICATION: US/10/522,592

TIME: 13:27:10

Input Set : N:\DA\pto.da.txt

Output Set: N:\CRF4\02222005\J522592.raw

3 <110> APPLICANT: MADJAR, Jean-Jacques
 4 BERTHOMME, Herv,
 6 <120> TITLE OF INVENTION: Novel method for analyzing nucleic acid and use thereof
 7 for evaluating the degree of mRNA editing of the
 8 serotonin 5-HT2C receptor
 10 <130> FILE REFERENCE: 347153
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/522,592
 C--> 12 <141> CURRENT FILING DATE: 2005-01-25
 12 <150> PRIOR APPLICATION NUMBER: PCT/FR 2003/002 339
 13 <151> PRIOR FILING DATE: 2003-07-24
 15 <150> PRIOR APPLICATION NUMBER: FR 02/09 524
 16 <151> PRIOR FILING DATE: 2002-07-26
 18 <160> NUMBER OF SEQ ID NOS: 37
 20 <170> SOFTWARE: PatentIn Ver. 2.1

pp1,3
 Does Not Comply
 Corrected Diskette Needed

ERRORED SEQUENCES

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 842 <211> LENGTH: 13
 E--> 843 <212> TYPE: ARN ? invalid The only valid <212> responses for
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 846 <220> FEATURE: RNA. FYI:
 847 <223> OTHER INFORMATION: Description of the artificial sequence: DNA derived
 848 from the mRNA coding for human 5-HT2C receptor
 850 <400> SEQUENCE: 33
 851 auacguaauc cua 13 use
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 874 from the mRNA coding for human 5-HT2C receptor

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/522,592

DATE: 02/22/2005

TIME: 13:27:10

Input Set : N:\DA\pto.da.txt

Output Set: N:\CRF4\02222005\J522592.raw

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901 <400> SEQUENCE: 35
902 canuncgunn uccunuu

17

SEQUENCES LISTING

<110> MADJAR, Jean-Jacques
BERTHOMME, Hervé

<120> Novel method for analyzing nucleic acid and use thereof
for evaluating the degree of mRNA editing of the
serotonin 5-HT₂ receptor

do not use subscripts

The types of error shown exist throughout
the Sequence Listing. Please check subsequent
sequences for similar errors.

The types of error shown exist throughout
the Sequence Listing. Please check subsequent
sequences for similar errors.

The CRT
software
cannot
process
them.

delete all foreign accent marks.
(or Scientific symbols)

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/522,592

DATE: 02/22/2005

TIME: 13:27:11

Input Set : N:\DA\pto.da.txt

Output Set: N:\CRF4\02222005\J522592.raw

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L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
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